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(cgn2_6/ptodata/2/iaa/5A_COMB.pep:*

(cgn2_6/ptodata/2/iaa/5B_COMB.pep:*

(cgn2_6/ptodata/2/iaa/6A_COMB.pep:*

(cgn2_6/ptodata/2/iaa/6B_COMB.pep:*

(cgn2_6/ptodata/2/iaa/6B_COMB.pep:*

(cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*

(cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.
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US-09-419-568F-28

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US-09-134-28-28

US-08-419-568F-27

US-08-418-27-16

US-08-418-27-16

US-08-418-28-27

US-08-418-28-28

US-08-418-28-28

US-08-170-113-2

US-08-170-113-2

US-08-452-613-2

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US-08-410-6548-1

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Listing first 45 summaries
                                          OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Sequence 2, Application US/09870574

Fatent No. 6551799

GENERAL INFORMATION

APPLICANT: Gurney, Austin L.

APPLICANT: Xie, Ming-Hong

APPLICANT: Xie, Ming-Hong

APPLICANT: Maruoka, Ellen M.

APPLICANT: Mod, William I.

TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDERS

TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDERS

TITLE OF INVENTION UNMER: US/09/870,574

CURRENT APPLICATION NUMBER: US 60/169,495

PRIOR PLILING DATE: 2000-05-22

PRIOR PLILING DATE: 2000-05-22

PRIOR FILING DATE: 2000-08-24

NUMBER OF SEQ ID NOS: 7

LENGTH: 179
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100.0%; Pred. No. 1.9e-100;
.ive 0; Mismatches 0;
            US-08-270-805C-4
US-08-410-6548-4
US-08-411-4
US-08-170-113-4
US-08-643-810A-4
US-08-542-624A-4
US-08-552-624A-4
US-08-452-624A-4
US-08-410-6548-3
US-08-410-6548-3
US-08-410-6548-3
US-08-170-113-3
US-08-170-113-3
US-08-170-113-3
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Best Local Similarity 100.
Matches 179; Conservative
                                   1147
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ORGANISM: Homo Sapien
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Sequence 17 Application US/09419568F

Fatent No. 6331613

GENERAL INFORMATION:
APPLICANT: Dumoutier. Laure
APPLICANT: Dumoutier. Laure
APPLICANT: Loubed, Jamila
APPLICANT: Loubed, Jean-Christophe
TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T
TITLE OF INVENTION: 180912.2
CURRENT APPLICATION NUMBER: US/09/419,568F
CURRENT PILING DATE: 1999-10-18
FRIOR FILING DATE: 1999-07-16
PRIOR FILING DATE: 1999-07-16
PRIOR FILING DATE: 1999-07-16
NUMBER OF SEQ ID NOS: 29
SEQ ID NOS: 29
                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Dumoutier, Laure
APPLICANT: Loubed, Jamila
APPLICANT: Loubed, Jamila
APPLICANT: Renauld, Jean-Christophe
TITLE OF INVENTION: ISolated Nucleic Acid Molecules which Encode T Cell Inducible Far
TITLE OF INVENTION: (TIFS)
TITLE OF INVENTION: (TIFS)
FILE REFERENCE: LUD 5543
CURRENT APPLICATION NUMBER: US/09/178,973B
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61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEILFPQSDRFRPYMQEVVP 120
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; Pred. No. 5.9e-79;
19; Mismatches 20
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; Pred. No. 5.9e-79;
19; Mismatches 20
                                                                                                                                                                                                       Sequence 15, Application US/09178973B
Patent No. 6274710
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SEQ ID NO 15
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Best Local Similarity 78.2%,
Matches 140; Conservative
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Best Local Similarity 78.23
Matches 140; Conservative
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US-09-419-568F-27
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US-09-419-568F-27
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APPLICANT: Louned, Jamila
APPLICANT: Louned, Jamila
APPLICANT: Louned, Jamila
TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible F9
TITLE OF INVENTION: TIPE)
TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE OF INVENTION NUMBER: US/09/354,243B
CURRENT APPLICATION NUMBER: US09/178,973
PRIOR FILING DATE: 1998-10-26
NUMBER OF SEQ ID NOS: 29
LENGTH: 179
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                                                            APPLICANT: Louded, January
APPLICANT: Louded, January
APPLICANT: Louded, January
APPLICANT: Louded, Jean-Christophe
TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Ce
TITLE OF INVENTION: (TIPS) The Proteins Encoded, and Uses Thereof
FILE REFERENCE: LUD 5543.2
CURRENT APPLICATION NUMBER: US/09/419,568F
CURRENT PILING DATE: 1999-10-18
PRIOR APPLICATION NUMBER: US09/154,243
PRIOR PILING DATE: 1999-10-16
PRIOR APPLICATION NUMBER: US09/178,973
PRIOR FILING DATE: 1998-10-26
NUMBER OF SEQ ID NOS: 29
SEQ ID NO 28
LENGTH: 179
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97.8%; Pred. No. 8.3e-98;
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Pred. No. 8.3e-98;
2; Mismatches 2
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Patent No. 6359117
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Best Local Similarity 97.8
Matches 175; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
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Matches 175; Conserv
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                                                                                                                                                                                                                                                                                 61 KEASLADNNTDVRLIGEKLFRGVSAKDQCYLMKQVLNFTLEDILLEPQSDRFRPYMGEVVP 120
                                                                                                                                                                      9
                                                                                                                                                                                         121 FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                            1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Di-Ewei Hsu
APPLICANT: Kevin K. Moore
APPLICANT: Hergen Spits
TITLE OF INVENTION: Use Of Interleukin10 in Adoptive Immunotherapy
TITLE OF INVENTION: Of Cancer
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                                                                         Length 179;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 170;
                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                       Score 720; DB 3;
Pred. No. 1.8e-78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12.1%; Score 109; DB 1; 26.7%; Pred. No. 3.7e-05; tive 24; Mismatches 68
                                                                  Query Match
Best Local Similarity 77.7%; Pred. No. 1.8e
Matches 139; Conservative 20; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
CITY: Kenliworth
STATE: New Jersey
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 07033
ZUP: 07033
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: 7.5.3
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/270,805C
FILING DATE: 05-Ualy-1994
CLASSIFICATION: 424
APPLICATION 1424
APPLICATION WINBER: US 07/995,564
FILING DATE: 23-Dec-1992
APPLICATION NUMBER: US 07/641,342
FILING DATE: 16-Jan-1991
APPLICATION NUMBER: US 07/641,342
FILING DATE: 16-Jan-1991
ATTORNEY/AGENT INFORMATION:
NAME: FOALKe, Cynthia L.
REGISTRATION NUMBER: 32,364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/08270805C Patent No. 5776451
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         908-298-2987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 170 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 26.77
Matches 43; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 170 amino acid
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ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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       ; OKCANAS:. ....
US-09-178-973B-16
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APPLICANT: Dumoutier, Laure
APPLICANT: Dumoutier, Laure
APPLICANT: Dumoutier, Laure
APPLICANT: Lounded, Jean-La
APPLICANT: Renauld, Jean-Christophe
TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa
TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE REFERENCE: LUD 5543.1
FILE REFERENCE: LUD 5543.1
CURRENT APPLICATION NUMBER: U5/09/354,243B
FRIOR PELICATION NUMBER: U5/09/354,243B
FRIOR PELICATION NUMBER: U5/09/178,973
FRIOR FILING DATE: 1998-10-26
NUMBER OF SEQ ID NOS: 29
LENGTH: 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Dumoutier, Laure
APPLICANT: Louhed, Jamia
APPLICANT: Louhed, Jamia
APPLICANT: Louhed, Jamia
TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
TITLE OF INVENTION: (TIFE)
TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
FILE REFERENCE: LUD 5543
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                                                                                                      61 KEASLADINITDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQFYMQEVVP 120
                                                                                                                                    KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
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                       60
                                             121 FLARLSNRLSTCHIEGDDLHIORNVOKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 MAVLQKSMSFSLMGTLAASCLLLIALMAQEANALPVNTRCKLEVSNFQOPYIVNRTFMLA
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80.2%; Score 724; DB 4; Length 179;
Best Local Similarity 78.2%; Pred. No. 5.9e-79;
Matches 140; Conservative 19; Mismatches 20; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/178,973B
CURRENT FILLING DATE: 1998-10-26
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                          Sequence 27, Application US/09354243B Patent No. 6359117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16, Application US/09178973B Patent No. 6274710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                             RESULT 6
US-09-354-243B-27
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US-09-178-973B-16
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LENGTH: 179
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73 RLIGEKLFHGVSMSERCYLMKOVLNFTLEEVLFPQSDRFQPYMQEVVPFLAR----LSNR 128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11 CLVLLYLAPECGG----TDQC----DNPPQMLRDLRDAFSRVKTFFQTKD-----EVDN 56
                                                                                                                                                                                                                                                                                                                        APPLICANT: Rene de Waal Malefyt
APPLICANT: Di-Hwei Hsu
APPLICANT: Di-Hwei Hsu
APPLICANT: Anneo O'Gatra
APPLICANT: Herneo O'Gatra
APPLICANT: Herneo O'Gatra
TITLE OF INVENTION: Use of An Interleukin-10 Antagonist to Treat
TITLE OF INVENTION: A B Cell Mediated Autoimmune Disorder
NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12.1%; Score 109; DB 2; Length 170; 26.7%; Pred. No. 3.7e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                              129 LSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDL 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/410,654
FILING DATE: 24-MAR-1995
APPLICATION NUMBER: US 08/229,854
FILING DATE: 19-APR-1994
APPLICATION NUMBER: US 07/926,853
FILING DATE: 06-AUG-1992
APPLICATION NUMBER: US 07/742,129
ATONNEY, AGENT INFORMATION:
NAME: FOLIKE, CYATCHA I.
REGISTRATION NUMBER: 32,364
REFERENCE/DOCKET NUMBER: 32,364
REFERENCE/DOCKET NUMBER: DX0221KQ1GD
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
CITY: Kenilworth
STATE: Redilanders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Microsoft Word 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/474,851
FILING DATE: 07-UN-1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                 Sequence 2, Application US/08474851
Patent No. 5837232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 908-298-5388
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 170 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: Macintosh
OPERATING SYSTEM: 7.5.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            43; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 07-JUN-CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 43; Conserv
                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07033
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                                                                                                                                           73 RLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVPFLAR----LSNR 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11 CLVLLYLAPECGG----IDQC---DNFPQMLRDLRDAFSRVKTFFQTKD-----EVDN 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 CLLLLALLVQGGAAAPISSHCRLDKSNFQQ-----PYITNRTFMLAKEASLADNNTDV 72
       20 CLILLALLVQGGAAAPISSHCRLDKSNFQQ-----PYIINRTFMLAKEASLADNNTDV 72
                                                       11 CLVLLYLAPECGG----TDQC----DNFPQMLRDLRDAFSRVKTFFQTKD-----BVDN 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        68; Indels 26; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rene de Waal Malefyt
APPLICANT: Rene de Waal Malefyt
APPLICANT: Di-Heei Ksu
APPLICANT: Di-Heei Ksu
APPLICANT: Hergen Spits
TITLE OF INVENTION: Use of Interleukin-10 to Treat
TITLE OF INVENTION: Septic Shock
NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
ADDRESSED: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
                                                                                                                                                                                                                                      116 LRRCHRFLPCENKSKAVBQIKMAFNKLQEKGIYKAMSEFDI 156
                                                                                                                                                                                                          129 LSTCHIEGDDLHIORNVOKLKDTVKKLGESGEIKAIGELDL 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA.
APPLICATION NUMBER: US 08/229,854
PILING DATE: 19-AR-1994
APPLICATION NUMBER: US 07/926,853
FILING DATE: 06-AUG-1992
APPLICATION NUMBER: US 07/742,129
FILING DATE: 06-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: POULKe, CYNTHIA I.
REGISTRATION NUMBER: 32,364
REFERNCE/POCKET NUMBER: DX0221KQ1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/410,654B
FILLIO DATE: 24-MAR-1995
CLASSIFCATION: 424
                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/08410654B
Patent No. 5833976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SYSTEM: 7.5.3
Microsoft Word 5.1a
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INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Floppy disk
Macintosh
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New Jersey
USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: Macinto OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
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73 RLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVPFLAR----LSNR 128
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                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Knappe, Andrea
APPLICANT: Fickenscher, Helmut
APPLICANT: Fickenscher, Helmut
APPLICANT: Fickenscher, Helmut
APPLICANT: Fickenscher, Bernhard
TITLE OF INVENTION: MAMMALIAN CYTOKINE; RELATED REAGENTS
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSE: DNAK Research Institute
STATE: 910 California Avenue
CITY: Palo Alto
STATE: California Avenue
CITY: Palo Alto
COMPUTER: Babbabe FORM:
MEDING TYPE: Floppy disk
COMPUTER: Exabbabe FORM:
MEDING TYPE: Floppy disk
COMPUTER: Date: Ploppy disk
COMPUTER: DATE: 22-SEP-1997
CLASSIFICATION NUMBER: US 60/027,368
FILING DATE: 23-SEP-1996
ATTORNEY CALING DATE: 23-SEP-1996
ATTORNEY CALING DATE: 23-SEP-1996
ATTORNEY CALING DATE: 33-SEP-1996
ATTORNEY CALING NUMBER: 34,090
REFERENCE/DOCKET NUMBER: DX0644K
TELECOMMUNICATION INDMERR: DX0644K
TELECOMMUNICATION INPORMATION:
TELEBROWN CALING
TELEBROWN CALING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 170;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
12.1%; Score 109; DB 2; Length 17
Best Local Similarity 26.7%; Pred. No. 3.7e-05;
Matches 43; Conservative 24; Mismatches 68; Indels
       116 LRRCHRFLPCENKSKAVEQIKNAFNKLQEKGIYKAMSEFDI 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  129 LSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDL 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              116 LRRCHRFLPCENKSKAVEQIXWAFNKLQEKGIYKAMSEFDI 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 CLLLLALLVQGGAAAPISSHCRLDKSNFQQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/08170113
Patent No. 6106823
GENERAL INPORMATION:
APPLICANT: Woere, Kevin W.
APPLICANT: Moore, Kevin W.
APPLICANT: de Waal Malefyt, Rene
                                                                                                                          Sequence 4, Application US/08934959
Patent No. 5989867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              , TOPOLOGY: not relevant
; MOLECULE TYPE: peptide
US-08-934-959-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 170 amino acids TYPE: amino acid STRANDEDNESS: not releve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 650-496-1200
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 170 amino acid
                                                                                                        US-08-934-959-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 13
US-08-170-113-2
       В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLILILIALLVQGGAAAPISSHCRLDKSNFQQ-----PYITNRTFMLAKEASLADNNTDV 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12.1%; Score 109; DB 2; Length 170; 26.7%; Pred. No. 3.7e-05; Live 24; Mismatches 68; Indels
                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Rene de Waal Malefyt
APPLICANT: Di-Hwel Hsu
APPLICANT: Ahre O'Garra
APPLICANT: Hergen Spits
ITILB OF INVENTION: Use of Interleukin-10 to Modulate
ITILE OF INVENTION: Inflammation or T-Cell Mediated
ITILE OF INVENTION: Immune Function
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schering-Plough Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   68; Indels
                              129 LSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDL 169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Schering-Plough Corporation
STREET: 2000 Galloping Hill Road
CITY: Remilworth
STATE: New Jersey
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Foulke, Cynthia L.
REGISTRATION NUMBER: 32,364
REFERNCE/DOCKET NUMBER: DX0221KQ1GC
TELECOMUNICATION INFORMATION:
TELEPHONE: 908-2987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OMPRATURE: Macintosh
OPERATURE: Microsoft Word 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/481,560
FILING DATE: 07-UN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/410,654
FILING DATE: 24-MAR-1995
APPLICATION NUMBER: US 08/229,854
FILING DATE: 19-APR-1994
APPLICATION NUMBER: US 07/926,853
FILING DATE: 06-AUG-1992
APPLICATION NUMBER: US 07/926,853
FILING DATE: 10-APR-1194
APPLICATION NUMBER: US 07/926,853
FILING DATE: 10-APR-1199
APPLICATION NUMBER: US 07/926,853
FILING DATE: NG-AUG-1991
ATTORNEY AGENT INFORMATION:
NUMBER: US 07/742,129
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                                                                                                                                                                     Sequence 2, Application US/08481560
Patent No. 5837293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 170 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 26.79
Matches 43; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-481-560-2
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ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C. STREET: 419 Seventh Street N.W., Suite 300

CITY: Washington STATE: D.C.

COUNTRY: U.S.A. ZIP: 20004

E: Floppy disk IBM PC compatible

COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

APPLICANT: GROWHOJ LARSEN, Christian APPLICANT: GESSER, Borbala TITLE OF INVENTION: IMMUNOMODULATORS NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS:

GENERAL INFORMATION: Patent No. 6159937

```
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/765,094C
FILING DATE: 06-JAN-1997
PRIOR APPLICATION DATE:
APPLICATION NUMBER: PCT/DK95/00227
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATE:
APPLICATION NUMBER: DK 0800/94
FILING DATE: 05-JUL-1994

**ATTORNEY/AGENT INFORMATION:
**ATTORN
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                                    APPLICANT: Fluckinger, Anne-Catherine
APPLICANT: Banchereau, Jacques
TITILE OF INVENTION: TREATMENT OF NEOPLASTIC DISEASE WITH
TITLE OF INVENTION: INTERLEUKIN-10
                                                                                                                                                                                                                                                                                                                                                                                                                          STALL.
COMPUTRY: USA
ZIP: 94304-1104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/170,113
FILING DATE: 17-DEC-1993
FILING DATE: 17-DEC-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRICE APPLICATION DATA

APPLICATION NUMBER: US 07/933,419
FILING DATE: 21-AUC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/091,333
FILING DATE: 12-UUL-1993
FILING DATE: 12-UUL-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/020,018
FILING DATE: 17-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/830,496
FILING DATE: 04-FEB-1992
PRIOR APPLICATION NUMBER: US 07/641,347
FILING DATE: 16-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAMB: CANDING Edwin P.
REGISTRATION NUMBER: 34,090
REGISTRATION NUMBER: 34,090
REGISTRATION NUMBER: 34,090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DX0116Q1KX
                                                                                                                                                                                                        NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: DNAX Research Institute
STREET: 901 California Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-852-9196
de Vries, Jan E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 415-496-1200
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       170 amino acids
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                               STREET: 901 Califo
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 43; Conserve
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NAME: COOPER, IVer P.
REGISTRATION NUMBER: 28,005
REGISTRATION NUMBER: GRONHOJ-LARSEN=1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)628-5197
TELEFAX: (202)737-3528

170 amino acids

single

TYPE: amino acid STRANDEDNESS: si

INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:

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73 RLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVPFLAR----LSNR 128
                                                                                                                                                                                                                                                     20 CLLLLALLVQGGAAAPISSHCRLDKSNFQQ-----PYITNRTFMLAKEASLADNNTDV
                                                                                                                    68, Indels 26;
                                                                                 Length 170;
                                                                                                                                                                                                                                                                                                      129 LSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDL 169
                                                                                                                                                                                                                                                                                                                              Sequence 26, Application US/09082797
Patent No. 6168791
GENERAL INFORMATION:
APPLICANT: GENSER, Borbala
TITLE OF INVENTION: INMUNOMODULATORS
NUMBER OF SEQUENCES: 27
CORRESPONDER ADDRESS:
ADDRESSE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 419 Seventh Street N.W., Suite 300
CITY: Washington
                                                                           Query Match
Best Local Similarity 26.7%; Pred. No. 3.7e-05; Matches 43; Conservative 24; Mismatches 68
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                   US-08-765-094C-26
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68; Indels

20 CLLLLALLVQGGAAAPISSHCRLDKSNFQQ-

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43; Conservative

RESULT 14 US-08-765-094C-26 ; Sequence 26, Application US/08765094C

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73 RLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVPFLAR----LSNR 128
                                                                     ----PYITNRTFMLAKEASLADNNTDV 72
                                                                                                                          129 LSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDL 169
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CORFERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/082,797
FILING ADDICATION DATA:
APPLICATION NUMBER: 08/765,094
FILING ADPLICATION DATA:
APPLICATION NUMBER: DR 0800/94
FILING ADDICATION DATA:
APPLICATION NUMBER: GR 0800/94
FILING ADDICATION INFORMATION:
TELEPRATION LIPORMATION:
TELEPRATION NUMBER: GR 080
TELEPAX: (202)737-3528
INPORMATION FOR SEQ ID NO: 26:
SEQUENCE CHRACTERISTICS:
LENGTH: 170 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: Amino acids
TYPE: AMINORASE. C. 10.70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: Single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-09-082-797-26
D.C.
U.S.A.
                                                                         20004
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73 RLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVPFLAR----LSNR 128 20 CLILLALLVQGGAAAPISSHCRLDKSNFQQ-----PYITNRTFWLAKEASLADNNTDV 72 68; Indels 26; Gaps Query Match 12.1%; Score 109; DB 3; Length 170; Best Local Similarity 26.7%; Pred. No. 3.7e-05; Matches 43; Conservative 24; Mismatches 68; Indels 129 LSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDL 169 ò

Search completed: December 5, 2003, 18:21:31 Job time : 29.64 secs

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Scoring table:

Searched:

Database

Perfect score:

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Sequence:

Human IL-TIF prote Mouse TIF beta pro Mouse interleukin-

ZCYTO18 pept ZCYTO18 pept IL-10. Synt interleukin-

Human Viral Human

Interleukin interleukin-interleukin-

Interleukin-

Viral Viral Viral Viral

Mouse TIF alpha pr Mouse T cell deriv

Mouse ZCYTO18 prot Murine T cell indu

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Human GIL-19 protein that shows a high degree of homology to IL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cell proliferation; immune stimulation; immune suppression; haematopoiesis regulation; tissue growth; inflammation; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; GIL-19/AE289; IL-10; interleukin-10; nutrition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Xuan
                                                                                                        AAU83713
ABU71582
ABU71582
ABU712067
ABU72185
ABU72185
ABU72185
ABU72185
ABU72185
AAU89091
AAES6559
AAY828777
AAU89090
                  ABB95599
AAU76909
                                                                     ABB84993
AAE19237
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AAE05052
AAY92878
AAE28614
AAE19236
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AAE05049
AAE05051
AAR26403
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AAR41665
AAR42643
AAW46586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human GIL-19/AE289 protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB36292 standard; Protein; 179 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Spaulding V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-APR-2000; 2000WO-US11479.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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Jacobs K, Fouser L,
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 Human T cell induc
Human cytokine, ZC
Human IL-TIF polyp
Human PKO10096. H
Amino acid sequenc
Human extracellula
Human interleukin-
Human interleukin-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human GIL-19/AE289
                                                                                                                 ; Search time 58.9323 Seconds (without alignments)
482.113 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                903
1 MAALQKSVSSFLMGTLATSC......EIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Description
             GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                    1107863 seqs, 158726573 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUMMARIES
                                                                                                                   5, 2003, 16:39:13
                                                                                                                                                                                                                                                                                                                                                                                                                                             Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                  OM protein - protein search, using sw model
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AAY92879
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AAB48074
AAE28608
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Gapop 10.0 , Gapext 0.5
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: /SIDS1/qcqdata/q
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Maximum DB seg length: 200000000
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                                                                                                                                                                                                                                                                                                                                1 MAALQKSVSFLMGTLATSCILLLALLVQGGAAAPISSHORLDKSNFQQPYITNRTFMLA 60
                                                                                                                                                                                                                                                                                                                                                                                                                            FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides the protein and coding sequences for the novel human GIL-19/A289 protein. The protein shows homology to interleukin-10 (IL-10) and is assumed to be a cytokinne. It can be used in the regulation of cell proliferation and differentiation, haematopoiseis, immune stimulation or suppression, tissue growth and tumour inhibition. In addition, it also has uses in the treatment of inflammation and in nutrition,
                                                                                                                                                                                                                                                                                                     1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This is the human T cell derived inducible factor (TIF). The gene was mapped to chromosome 12415. The human TIF was identified based on homology to a murine TIF, which was identified by subtraction cloning
(interleukin)-10, useful in upregulation of humoral immune responses,
                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid molecule encoding a T cell derived inducible factor
for treating asthma, an allergy or lymphoma
                   as an antiinflammatory agent and as a modulator of immune responses associated with injury -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LIF; T cell derived inducible factor; interleukin 9; STAT; IL-9;
Anti-asthmatic; anti-allergic; cytostatic; inhibitor; antagonist;
                                                                                                                                                                                                                                                                               ;
0
                                                                                                                                                                                                                                                 Length 179;
                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                          100.0%; Score 903; DB 21;
100.0%; Pred. No. 2.7e-86;
iive 0; Mismatches 0;
                                                            Claim 9; Page 59-60; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY92879 standard; Protein; 179 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 26; Fig 1; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (LUDW-) LUDWIG INST CANCER RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human T cell inducible factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98US-0178973.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99WO-US24424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-422495/36.
N-PSDB; AAA28839, AAA28840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                          Local Similarity 100.
1es 179; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dumoutier L, Louhed J,
                                                                                                                                                                                                               179 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chromosome 12q15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WC200024758-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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16-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-SEP-2000
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                                                                                                                                                                                                               Sequence
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interleukin 9 (IL-9). BW5147, can be grown in vitro, without the need to add any cytokines to its culture medium. Many IL-9 activities are mediated by activation of STAT transcription factors. The novel TIFs were STAT activation in CFTAT transcription factors. The novel TIFs were STAT activation in cells. They can be used, e.g. in the stimulation of targeted tissues. Their inhibitors or antagonists can be used to retard, prevent or inhibit differentiation of targeted tissues. Their inhibitors of attament of sathmalliers and their coding sequences are useful in the treatment of asthmalliers compounds that inhibit or activate T cell induced factor activity in a cell (claimed).
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                                                                                                                                                                                                                                                                                                                                                                                                                              FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFWSLRNACI 179
                                                                                                                                                                                                                                                                                                                                                                                                            121 FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                                                                                                                                                                      1 MAALQKSVSSFIMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                                                                                                                                                                                                                                                   61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP
                                                                                                                                                                                                                                                                                           1 MAALOKSVSSFLMGTLATSCLLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel human cytokine polypeptide, ZCYTO18, useful for treating cancer
                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, cytostatic, cytokine, ZCYTO18 protein, genetic abnormality, cancer, inflammation, gene therapy.
                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                     Length 179;
                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                               100.0%; Score 903; DB 21;
100.0%; Pred. No. 2.7e-86;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1..33
/label= Signal_peptide
34..179
/label= Mature_ZCYTO18_protein
145..150
/label= Hydrophilic_region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human cytokine, ZCYTO18 protein #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAE04538 standard; Protein; 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-DEC-1999; 99US-0471767.
01-DEC-2000; 2000US-0250841.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Presnell SR, Kindsvogel W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                            Matches 179; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-408648/43.
                                                                                                                                                                                                     179 AA;
                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; AAD09719
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                                                                                                                                                                                                      Seguence
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Peptide
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Length 179; Indels 9 9

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IL-TIF induced or IL-9 induced inflammation, and suppressing an inflammatory response in a mammal with inflammation. Heteromeric/multimeric receptor polypeptides such as soluble zcytor 16/CRE2-4 can be used to reduce progression and symptoms of cancer. Zcytor16 polypeptides can also be used to detect IL-TIF levels which is indicative of arthritis) and cancer. Antibodies inflammatory states (e.g. rheumatoid arthritis) and cancer. Antibodies that bind zcytor16 polypeptides and the polypeptides are useful for the treatment of inflammation, inflammatory diseases (e.g. infection, asthma, inflammatory bowel diseases, rheumatoid arthritis and atherosolerosis) and autoimmune diseases. The antibodies and zcytor16 polymucleotides are also useful for detecting cancer. The present sequence represents the human IL-TIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 KEASLADINITDVRLIGEKLFHGVSMSERCYLMKQVINPTLEEVLFPQSDRFQPYNQEVVP 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121 FLARLSNRLSTCHIEGDDLHIQRNVQXLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 MAALQKSVSSFIMGTLATSCILLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 KEASLADNNTDVRLIGEKLFHGVSKSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
        differentiation of hematopoietic cell(s) (progenitors); reducing
                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 903; DB 22;
100.0%; Pred. No. 2.7e-86;
cive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB87602 standard; Protein; 179 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; PRO protein; mapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US21090
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2000US-0199397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-MAY-2000; 2000WO-US14042.
05-JUN-2000; 2000US-0209832.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 179; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                           179 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PRO10096
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18-FEB-2000;
18-FEB-2000;
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01-MAR-2000;
03-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-SEP-1999;
15-SEP-1999;
07-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-MAR-2001
                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB87602;
                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMGEVVP 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
                                                                                                                                                                                                                                                                                                                                                                                        9
                                                                                                                                                                                                                                                                                                                                                                                                       FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a human cytokine receptor polypeptide, designated zcytor16. The zcytor16 polypeptide can be expressed by standard recombinant methodology and can bind to IL-TIF (undefined). The zcytor16 protein is useful for: inhibiting IL-TIF induced proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New soluble cytokine receptor polypeptides and polynucleotides, useful for diagnosing and treating cancer and inflammatory conditions -
                                                                                                                                                                                                                                                                                                                                                                             1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                     The patent discloses novel human cytokine, ZCYTO18 protein and its corresponding DNA. ZCYTO18 protein induces proliferation of cells expressing zcytor11, a receptor for SCYTO18 or induces cytochaticty in K5626 cells. ZCYTO18 DNA is useful for detecting a genetic abnormality in a patient. ZCYTO18 DNA and its antibodies are useful for detecting cancer and inflammation. ZCYTO18 protein is useful for killing cancer cells. It is useful for increasing platelets in a patient or injured tissue. It is also used in gene therapy. The present sequence is novel human cytokine, ZCYTO18 protein.
                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytokine receptor; zcytor16; IL-TIF; antiinflammatory; cytostatic; antirheumatic; antiarthritic; antiasthmatic; antiatherosclerotic; immunosuppressive; chromosome 6q24.1-25.2; human.
                                                                                                                                                                                                                                                                                                                                             .
0
                                                                                                                                                                                                                                                                                             100.0%; Score 903; DB 22; Length 179; 100.0%; Pred. No. 2.7e-86; ive 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 17; Page 195-196; 210pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB62664 standard; Protein; 179 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kindsvogel W,
Claim 4; Page 142; 167pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-DEC-1999; 99US-0169049.
13-SEP-2000; 2000US-0232219.
31-OCT-2000; 2000US-0244610.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-DEC-2000; 2000WO-US32703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                      Matches 179; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human IL-TIF polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Xu W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-356158/37
                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                     179 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; AAF83741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200140467-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUN-2001.
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                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121
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                                                                                                                                                                                                                                                                                               Query Match
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RESULT 4 AAB62664

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Godowski PJ;

Goddard A, Wood WI;

Gerritsen ME, Watanabe CK,

Filvaroff E, 7, Gurney AL,

Eaton DL, Fi Grimaldi CJ,

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22-MAY-2000; 2000WO-US14042

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The present sequence is a human PRO polypeptide (secreted and transmembrane). The PRO protein, and PRO agonists, PRO antagonists or anti-PRO antibodies are useful for preparation of a medicament useful in the treatment of a condition which is responsive to the PRO protein, agonists, antagonists or anti-PRO antibodies. The PRO protein may also be employed as molecular weight markers for protein electrophoresis. The PRO coding sequence has applications in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, secreted protein, transmembrane protein, PRO196; PRO444; PRO183; PRO185; PRO216; PRO215; PRO217; PRO217; PRO217; PRO517; PRO4170; PRO5183; PRO1308; PRO183; PRO183; PRO183; PRO5189; PRO5189; PRO5180; PRO5940; PRO533; PRO318; PRO5180; PRO5180; PRO5180; PRO5181; PRO4181; PRO4181; PRO4180; PRO5180; PRO518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 KEASLADINITDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNPQQPYITNRTFMLA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           121 FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRO6309; cell death; genetic disorder; transgenic animal; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                          Eighty four nucleic acids encoding PRO polypeptides, useful in molecular biology, including use as hybridization probes, and in chromosome and gene mapping.
                                                                                                                                                                                                                                                                                                                                                                                                       Length 179;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amino acid sequence of human polypeptide PRO10096.
                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 903; DB 22;
100.0%; Pred. No. 2.7e-86;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "N-myristoylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "N-glycosylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "N-glycosylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1..33
/note= "signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB31210 standard; Protein; 179 AA
                                                                                                                                       Claim 12; Fig 154; 278pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                         179; Conservative
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/note=
68..72
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  WPI; 2001-183260/18.
                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                            179 AA;
                      N-PSDB; AAF92134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified-site
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                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                        Best.Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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g
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'note= "N-myristoylation site"

/note= "N-glycosylation site"

WO200077037-A2

21-DEC-2000.

Modified-site

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The present sequence represents a human secreted and transmembrane polypeptide. The specification describes human polypeptides, designated PRO196, PRO444, PRO183, PRO116, PRO215, PRO217, PRO242, PRO288, PRO186, PRO186, PRO187, PRO1872, PRO187, PRO246, PRO187, PR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVINFTLEEVLFPQSDRFQPYMQEVVP 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated nucleic acid molecule encoding a PRO polypeptide which is a transmembrane polypeptide is useful for gene therapy and identification of related polypeptides -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 MAALQKSVSSFIMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Baker KP, Botstein DA, Desnoyers L, Eaton DL;
ng S, Gao W, Gerber H, Gerritsen ME, Goddard A;
Gurney AL, Kljavin IJ, Mather JP, Napier MA, Pan J;
MA, Stewart TA, Tumas D, Watanabe CK, Williams PM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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100.0%; Pred. No. 2.7e-86;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 12; Fig 64; 244pp; English.
                                                                                                                99WO-US21090
                                                                                                                                                                                         99WO-US28313
                                                                                                                                                                                                                                                               99WO-US28565
                                                                                                                                                                                                                                                                                                                                    2000WO-US00219
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                                                                                                                                                                                                                                                                                                                                                                                                              2000WO-US04342
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                                                                         99WO-US20594
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Best Local Similarity 100.
Matches 179; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ashkenazi AJ, Baker Ki
Ferrara N, Fong S, G;
Godowski PJ, Gurney A!
Paoni NF, Roy MA, St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC.
                                      99WO-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-050091/06.
N-PSDB; AAC87053.
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Paoni NF, Roy Fr.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    179 AA;
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18-FEB-2000;
22-FEB-2000;
                                                                                                                                                                             30-NOV-1999;
01-DEC-1999;
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17-MAY-2000;
                                                                                                                                                  5-SEP-1999;
                                                                                                                                                                                                                                                           02-DEC-1999;
07-DEC-1999;
                                                                                                                                                                                                                                                                                                                                    05-JAN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-MAR-2000;
                                          01-SEP-1999;
                                                                                                    15-SEP-1999;
17-AUG-1999
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FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI

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Extracellular signaling molecule, EXCS, anti-inflammatory, human; immunosuppressive; cytostatic, neuroprotective; gastrointestinal; virucide; antibacterial; anti-HIV; human immunodeficancy virus; antilifertility; cerebroprotective; noctropic; antiulcer; antifumgal; anticonvulsant; tranquilizer; neuroleptic; vasotropic; gynecological;
Human extracellular signaling molecule (EXCS) (ID 5571181CD1).
                                                                                                                                                  keratolytic; protozoacide; gene therapy.
                                                                                                                                                                                                                                                                                          l P, Burford N,
Patterson C;
                                             AAB48074 standard; protein; 179 AA
                                                                                                                                                                                                                                          99US-0144270.
                                                                                                                                                                                                                   19-MAY-2000; 2000WO-US13975
                                                                                                                                                                                                                                                          99US-0157508
                                                                                                                                                                                                                                                                          (INCY-) INCYTE GENOMICS INC
                                                                            (first entry)
                                                                                                                                                                                                                                                                                           Lal P,
                                                                                                                                                                                                                                                                                                                WPI; 2001-025021/03.
                                                                                                                                                                                                                                                                                                 Lu DAM,
                                                                                                                                                                                                                                                                                          Yue H,
                                                                                                                                                                                                                                                                                                                        N-PSDB; AAC84310
                                                                                                                                                                                   WO200070049-A2
                                                                                                                                                                                                                                  19-MAY-1999;
15-JUL-1999;
30-JUL-1999;
                                                                                                                                                                   Homo sapiens
                                                                            19-MAR-2001
                                                                                                                                                                                                                                                          04-OCT-1999;
                                                                                                                                                                                                   23-NOV-2000
                                                                                                                                                                                                                                                                                                 Azimzai Y,
                                                            AAB48074;
                                                                                                                                                                                                                                                                                          Tang YT,
                                      AAB48074
                              RESULT
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New human extracellular signaling nucleic acids and polypeptides useful for diagnosing, treating and preventing infections and gastrointestinal, neurological, reproductive, and autoimmune/inflammatory disorders Claim 1; Page 94; 114pp; English.

The invention provides human extracellular signaling molecules (EXCS)
and polynucleotides which identify and encode EXCS. EXCS can be
CC expressed by standard recombinant methodology. The amino acid and nucleic
acid sequences of EXCS are useful for diagnosing, treating and
acid sequences of EXCS are useful for diagnosing, treating and
parceatitis), neurological (e.g. epilepsy, ischemic cerebrovascular
cd iscase, stroke), reproductive (infertility, ovulatory defects,
cdiscase, stroke), reproductive (infertility, ovulatory defects,
cdiscase, stroke), autoimmune /inflammatory deficiens, acquired
immunodeficiency syndrome (AIDS), Addison's discase), and cell
proliferative disorders including cancers (of the breast, adrenal gland,
bone). They may also be used to treat fatal familial insomnia,
cc nutritional and metabolic discases of the nervous system, myopathies,
cutritional and metabolic discases of the nervous system, myopathies,
cutridional and metabolic discases of the nervous system, myopathies,
cutridional and metabolic discases of the nervous system, sinfections
caused by parasites (malaria, leishmania, trypanosoma), viral
caterphyles) agents. The nucleic acids, polypeptides, antagonists,
cd agonists, pharmaceutical compositions, and antibodies may also be used
cf for treating or preventing disorders associated with increased or
decreased expression or activity of EXCS. EXCS polypucleds may also
be used to detect and quantify gene expression in biopsied tissues in
which expression of EXCS may be correlated with the disease, to determine

ò levels during therapeutic intervention, to detect the presence of associated disorders, as targets in microarray, to generate hybridization probes, and to detect differences in gene sequences among normal, carrier or affected individuals. Antibodies may also be used in diagnosing disorders, in monitoring patients being treated with EXCS agonists, antagonists or inhibitors. Sequences AAB48057-B48082 represent the EXCS 61 KEASLADNNTDVRLIGEKLFHGVSMSERCXLAKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120 1 MAALQKSVSSFLMGTIATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITVRTFMLA 60 121 FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179 1 MAALOKSVSSFIMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA 61 KEASLADNNIDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP expression of EXCS, to monitor regulation of EXCS 0; Length 179; Indels Score 903; DB 22; Pred. No. 2.7e-86; 0; Mismatches 100.0%; Conservative Similarity 179 AA; of the invention Matches 179; Sequence Query Match Best Local ð 셤 à 셤

AAE28608 standard; Protein; 179 RESULT 8 

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(first entry) 27-DEC-2002

Human IL-TIF protein #1.

Baughn MR;

Bandman O,

Cytokine receptor; Zcytorl6; IL-TIF; autoimmune disease; dermatological; difflammatory disease; inflammatory bowel disease; rheumatoid arthritis; asthma; systemic lupus erythematoous; mysathenia gravis; pancreatitis; diabetes; atherosalezosis; glomerulonephritis; gene therapy; cytostatic; immunosuppressive; nephrotropic; allergy; placental health; abortion; cancer; human

Homo sapiens.

WO200270655-A2.

12-SEP-2002

04-MAR-2002; 2002WO-US06267.

02-MAR-2001; 2001US-273035P. 27-MAR-2001; 2001US-279232P.

(ZYMO ) ZYMOGENETICS

Chen Kindsvogel W, Xu W, Presnell SR,

WPI; 2002-698750/75

N-PSDB; AAD45964.

New Zcytor16 polypeptide useful for treating autoimmune or inflammatory diseases, e.g. inflammatory bowel disease, rheumatoid arthritis, asthma, atherosclerosis, cancer or diabetes, or in assessing therapeutic aspects of IL-TIF -

Disclosure; Page 197-198; 221pp; English.

The invention relates to cytokine receptor designated as mouse Zcytor16 which can bind and antagonise the IL-TIF. The Zcytor16 polypeptide is useful in modulating the immune system by binding Zcytor16 ligand, and

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120
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                                                                                                                                                                                                                                                                                                                                                         1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAFISSHCRLDKSNFQQPYITNRTFMLA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                           121 FLARLSHRISTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor. It is useful for studying human inflammation or immune function, or for treating autoimmune or inflammation or immune inflammatory diseases such as inflammatory bowel disease, rhematoid arthritis, asthma, systemic lupus erythematosus, myasthenia gravis or alleray, atheroselerosis, cancer, diabetes, glomerulonephritis or pancreatitis, or in assessing therapeutic aspects of IL-TIF, chemical therapeutics, anti-IL-TIF antibodies, anti-Zcytor16 antibodies or Zcytor16 soluble receptors. Zcytor16 bNA and the anti-mouse Zcytor16 antibody are useful as probes in detecting gene expression and gene structure, such as in the diagnosis and/or prevention of spontaneous abortions or in monitoring placental health and function. It is also used in gene therapy. The present sequence is human IL-TIF protein.
                                                                                                                                                                                                                                                                                                                            1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                                                                                                                                                                                                                                                                                             61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLBEVLFPQSDRFQPYMQEVVP
                                                                                                                                                                                                                                                                                                   Gaps
preventing the binding of the ligand with endogenous Zcytor16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Interleukin-22; IL-22; human; cytostatic; antiinflammatory; antibacterial; virucide; fungicide; osteopathic; vulnerary; neuroprotective; immunosuppressive; antiarthritic; antidiabetic; antipsoriatic; antiarteriosclerotic; anticholesterolemic; antiallergic; cytokine; adjuvant.
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0
                                                                                                                                                                                                                                                                   Length 179;
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note= "IL-22 receptor binding motif"
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                                                                                                                                                                                                                                                               Score 903; DB 23;
Pred. No. 2.7e-86;
0; Mismatches 0;
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/label= Mature protein
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/label= Signal_peptide
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                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0%;
Matches 179; Conservative 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-FEB-2002; 2002WO-US05684
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human interleukin-22.
                                                                                                                                                                                                                                179 AA;
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                                                                                                                                                                                                                                Seguence
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61 KEASLADINITDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
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                                                                                                                                                                                                                                                The present sequence is the protein sequence of human interleukin-22 (IL-22), a cytokine involved in acute phase cresponses. The invention provides inhibitors of IL-22, especially an antibody which immunologically reacts with an IL-22 protein, antibody, or other IL-22 modulator, is useful for treating a pathological condition in need of IL-22 modulation, or associated with IL-22 activity in a subject, particularly a human. In particular, these conditions include septicaemia, autoimmune disorders (e.g. arthritis or rheumatoid arthritis, osteoarthritis, multiple sclerosis, myasthenia gravis, inflammatory bowel disease, infection by bacteria, viruses, parasites or fungi), or the regulation of inflammation and acute phase responses (e.g. unfaction by bacteria, viruses, parasites or fungi), or the regulation of inflammation and acute phase responses (e.g. wound healing process, cholesterol metabolism, oxygen free radical injury, activity as a realso useful for treating inflammatory pathology in the kidney (which is a result of necrosis due to ischemia) cor cancer (e.g. renal cell carcinoma), or for remodelling kidney tissue (all claimed). A subject's immune response to an antigen can be enhanced by administering the antigen and an embanced immunogenicity-augmenting amount of IL-22.
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                                                                                                                 New antibodies that immunologically react with an interleukin 22 (IL 22) protein, useful for treating conditions associated with IL-22 activity in humans, e.g. septicemia, multiple sclerosis, ischemia, atherosclerosis, allergies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 903; DB 23; Length 179; 100.0%; Pred. No. 2.7e-86;
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                        Xuan
                      Spaulding V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted/transmembrane protein PRO10096.
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                        'n
                                                                                                                                                                                                                  Claim 6; Page 10; 76pp; English.
                        Fouser
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                      Jacobs K, Pittman D,
                                                         WPI; 2002-698660/75
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                                                                              N-PSDB; ABQ81260
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Claim 20; Fig 154; 399pp; English.
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98US-089514P.
98WO-US19330.
99WO-US05028.
99WO-US10733.
                            97US-063435P.
97US-064215P.
98US-082797P.
98US-083495P.
                                                      98US-085579P.
98US-088811P.
98US-088824P.
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2000WO-US15264.
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2000WO-US04342.
                                                                                                                                                                                                        2000WO-US23328.
2000WO-US30873.
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28-FEB-2001; 2001WO-US06520.
20-JUN-2001; 2001WO-US19692.
              06-DEC-2001; 2001US-0006867
                                                                                                                                                                                                2000WO-US23522.
                                                                                                                                                                                                                     2000WO-US32378
                                                                                                                                                                                                                                                       09-JUL-2001; 2001WO-US21735
                                                                                                                                                                                                                                                                                  Baton DL, Filvaroff E,
Grimaldi JC, Gurney AL,
                                                                                                                                                                                                                                                                     (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                         WPI; 2002-731348/79.
                                                                                                                                                                                                                                                                                                                N-PSDB; ABS74454.
                                                                                                                                                                            30-MAR-2000; 22-MAY-2000; 202-JUN-2000; 2
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                                                                                                                                  01-SEP-1999;
                                                                                                                                                15-SEP-1999;
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29-AUG-2002
                                                        15-MAY-1998
                                                                    10-JUN-1998
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                                                                                                                     14-MAY-1999
CCCCCCCCCCCCCCCCCCCCCX8X4F4F4XBAX4F4F4XBXXBXAF4F4F4F4F4F4F4F4F4F4F4F4F4F4F4F4F4F
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Godowski PJ; Goddard A, Wood WI; Gerritsen ME, Watanabe CK, New isolated secreted and transmembrane PRO polypeptide useful for modulating biological activity of a cell, or for treating sports-related joint problems, osteoarthritis or rheumatoid arthritis

The invention relates to an isolated secreted and transmembrane PRO polypeptide having 80 % sequence identity to a sequence appearing as ABG95814 advantages or their associated signal peptide, or a sequence of an extracellular domain of the proteins with their associated signal peptide or lacking its associated signal peptide. Also included are the muteler acids encoding the proteins, vectors, host cells.

CC the nucleic acids encoding the proteins, vectors, host cells.

The proteins are useful for detecting a polypeptide designated as A, B, C or D polypeptide or D in a sample suspected of containing an A, B, C or D polypeptide, C or D in a sample watch a polypeptide designated as B, P, C or D y contacting the sample with a polypeptide designated as B, P, B, C or D in the sample where A is a PRO10272 polypeptide, B is a PRO2010 C or D/I polypeptide conjugate is indicative of the presence of an A, B, C or D polypeptide conjugate is indicative of the presence of an A, B, C or D polypeptide C polypeptide, H is a PRO10313 polypeptide, D a PRO10376 polypeptide, E is a PRO20310 C polypeptide, H is a PRO20313 polypeptide and I is a PRO1030 C polypeptide. The sample comprises a cell suspected of expressing the A, C B, C D polypeptide. The B, F is G, H or I polypeptide is labeled with a detectable label or is attached to a cell expressing a cell expressing a processing and useful for linking a bloactive molecule to a cell expressing a

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polypeptide designated as A, B, C or D or B, F, G, H or I. The bioactive molecule is a toxin, a radiolabel or an antibody. The bioactive molecule against them are useful for modulating a biological activity of a cell against them are useful for modulating a biological activity of a cell expressing a polypeptide designated as A, B, C or D or E, F, G, H, or I. The cell is killed. The proteins are useful for identifying agonists or antagonists, for the proteins are useful for identifying the treatment of a condition which is responsive to the proteins, as molecular weight markers for protein electrophoresis purposes, and as therapeutic agents for treating sports-related joint problems, as therapeutic agents for treating sports-related joint problems, and in chromosome and gene mapping, in the generation of anti-sense RNA and DNA, for the preparation of the proteins, to generate transgenic or knockout animals which are useful in the development and screening of therapy. The antibody is useful as a therapeutic agent, and in gene therapy. The antibody is useful as a therapeutic agent, in a diagnostic assay and for affinity purification of the protein from recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 KEASLADNNTDVRLIGEKLFHGVSMSERCYIMKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell culture natural sources. The present sequence represents a novel secreted or transmembrane protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
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100.0%; Pred. No. 2.7e-86;
tive 0; Mismatches 0; Indels 0
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Matches 179; Conservative
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ABB95599 standard; Protein; 179 AA. ABB95599

RESULT 11

ABB95599;

(first entry) 19-JUL-2002

Human; angiogenesis; PRO protein; cardiovascularisation; wound; cancer; theroscalerosis; cardiac hypertrophy; gene therapy; endothelial disorder; cardiant; cytostatic; antiangiogenic; hypotensive; vulnerary; antiarteriosclerotic. Human angiogenesis related protein PRO10096 SEQ ID NO: 354.

Homo sapiens.

WO200208284-A2

31-JAN-2002

09-JUL-2001; 2001WO-US21735

2000US-220624P 2000WO-US20710 2000US-222695P 25-JUL-2000; 28-JUL-2000;

2000US-0643657 2000WO-US23522 17-AUG-2000; 23-AUG-2000; 24-AUG-2000; HANDER STANDER STANDER

2000WO-US23328. 2000US-230978P. 2000US-000000P. 07-SEP-2000; 15-SEP-2000; ω

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The present invention provides the protein and coding sequences of human PRO proteins. These are useful for treating or diagnosing a cardiovascular, endothelial or angiogenic disorder, including cardiac hypertrophy, trauma, cancer, age-related macular degeneration, atherosclerosis, hypertension, arterial restenosis, rheumatoid arthritis, angina, myocardial infarctions, thrombophlebitis, lymphangitis, tumour angiogenesis (such as breast carcinoma and liver carcinoma) and wound healing. The present sequence is a PRO protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      One hundred and eighty seven nucleic acids encoding PRO polypeptides, useful in diagnosis and treatment of cardiovascular (e.g. myocardial infarction), endothelial or angiogenic disorders in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gerber H, Gerritsen ME, Goddard A;
L, Hillan KJ, Marsters SA, Pan J,
CK, Williams PM, Wood WI, Ye W;
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                                                                                                                                                                             2001US-0828366.
2001US-0854208.
2001US-0866028.
2001US-0866034.
2001US-0866034.
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2001US-0796498.
2001WO-US06520.
2001WO-US06666.
                                                                                                                                             2001US-0802706.
2001US-0808689.
2001US-0816744.
                                                                   2000WO-US32678
2000US-0747259.
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2001WO-US17800
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F, Watanabe CK,
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GURNEY A L.
HILLAN K J.
MARSTERS S A.
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GERRITSEN M E.
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STEPHAN J F.
WATANABE C K.
WILLIAMS P M.
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FERRARA N.
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28-FEB-2001; 2
28-FEB-2001; 2
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18-SEP-2000;
18-SEP-2000;
24-OCT-2000;
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20-DEC-2000;
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25-MAY-2001;
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KEASLADNNTDVRLIGEKLFHGVSMSERCYTAKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
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                                                                                                                                                         FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                              This invention relates to the protein and cDNA sequences of a novel soluble cytokine receptor polypeptide designated zcytorII, which binds interleukin-T-cell inducible factor (IL-TIF) or antagonises IL-TIF activity. The protein of the invention is useful for reducing IL-TIF. or IL-9 induced inflammation, and inhibiting IL-TIF-induced proliferation. The protein is also useful for suppressing an immune response in a mammal exposed to an antigen or pathogen. Soluble zcytorII receptor or heterodimeric polypeptide is useful for enhancing the in vivo killing of target tissues by directly stimulating a zcytorII receptor-modulated apoptotic pathway. IL-TIF is involved in promoting Th-type immune responses and antagonists of IL-TIF have beneficial use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            against diseases involving such immune responses. Soluble zcytorII heterodimers are useful as antagonists in inflammatory and immune diseases or conditions such as pancreatitis, type I diabetes (IDDM), pancreatic cancer, Graves disease, inflammatory bowel disease (IBD), Crohn's disease, colon and intestinal cancer, diverticulosis, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New soluble cytokine receptor which binds interleukin-T-cell inducible factor and antagonizes its activity in inflammatory and immune diseases such as cancer, diabetes, asthma, sepsis, psoriasis and autoimmune
                         1 MAALQKSVSSFIMGTLATSCLLLIALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
MAALQKSVSSFIMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
                                                                                                                     61 KEASLADINITDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Z-Cytor II; human, cytokine receptor; atopy; psoriasis; interleukin-T-cell inducible factor; II-TIF; allergy; asthma; receptor-modulated apoptosis; Thi; immune response; pancreatitis; type I diabetes; IDDN; pancreatic cancer; Graves disease; SIE; inflammatory bowel disease; IBD; Crohn's disease; colon cancer; inflammatory disease; diverticulosis; autoimmune disease; sepsis; multiple sclerosis; MS; systemic lupus erythematosus; myasthaenia gravis; rheumatoid arthritis; kidney dysfunction.
                                                                                                                                                                                                                                                                                                                                                                                                                        Human interleukin-T-cell inducible factor (IL-TIF).
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                                                                                                                                                                                                                                                                                                  AAU76909 standard; Protein; 179
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                                                                                                                                                                                                                                                                                                                                                                                  05-JUN-2002 (first entry)
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Paoni NF;

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Gaps

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100.0%; Score 903; DB 23; Length 179; 100.0%; Pred. No. 2.7e-86; cive 0; Mismatches 0; Indels 0

Query Match 100. Best Local Similarity 100. Matches 179; Conservative

us-10-084-298-2.rag

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Interleukin 22; IL-22; cytostatic; antiinflammatory; IL-22 antagonist; immunotherapy; PAPI; pancreatitis associated protein; receptor; IL-22R; IL-10Rbeta; bioactive molecule linkage; cell death; pancreatitis; pancreatic carcinoma; acinar cell carcinoma; human; mixed cell population pancreatic carcinoma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     erythematosus (SLE), myasthaenia gravis, rhojn zhoumatoid arthritis and IBD), sepsis, asthma, allergy and other atopic diseases, psoriasis and kidney dysfunction. Soluble zcytoriI receptor or heterodimeric receptor polypeptides are useful in vivo or in diagnostic applications to detect IL-TIE expressing cancers in vivo or in tissue samples and to prepare antibodies. ZcytoRII serves as a target for MAb therapy of cancer where an attagonising MAb inhibits cancer growth and targets immune-mediated killing. The present sequence represents the interleukin-T-cell inducible factor (IL-TIE) protein, the activity of this protein is inhibited by the ZcytoRII protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                121 FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                                                                                                                                                                                                                                                                                  1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
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disease (e.g. IDDM, multiple sclerosis (MS), systemic lupus
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/label= Mature_human_interleukin_22_(IL_22)
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/note= "Asn is N-myristolated"
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                                                                                                                                                                                                                                                                  0; Mismatches
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/label= Signal_peptide
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                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                179 AA;
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24-AUG-2000; 2000WO-US23328

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The present invention relates to a new polypeptide having at least 80% identity to a 101 amino acid interleukin (IL)-22 sequence. The invention cis useful for detecting IL-22x (IL-22 receptor) or IL-10xbeta polypeptide in a sample which involves contacting sample with an IL-22 polypeptide can determining the formation of an IL-22x/IL-22 polypeptide conjugate or an IL-10xbeta/IL-22 polypeptide conjugate. Preferably, the IL-22 polypeptide or an IL-10xbeta/IL-22 polypeptide conjugate. Preferably, the IL-22 polypeptide or an IL-10xbeta/IL-22 polypeptide or antibody that causes the death of the cell, to support. The polypeptide is also useful for linking a bioactive molecule. Cc of the apprehensive molecule and allowing binding of the IL-22 polypeptide with cinvolves contacting the cell with IL-22 polypeptide with convolves contacting the cell with IL-22 polypeptide with convolves contacting the orlypeptide thus linking the bioactive molecules to the cell. The molecules of the invention can also be used for modulating biological activity of cell expressing IL-22R or IL-10xbeta polypeptide. Cc the cell is the invention of PAPI (pancreating a ssociated protein) by pancreatic cells. The antibody is also useful for treating a pancreatic disorder such as acute or chronic pancreatitis, pancreatic carcinoma and for reducing the activated or inflamed condition carcinoma and for reducing the activated or inflamed condition contaction and pancreatic astonaction the pancreatic associated the human interleukin 22 (IL-22) protein of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; anglogenesis; cardiant; cytostatic; antiangiogenic; hypotensive; vulnerary; antiarteriosclerotic; PRO agonist; PRO antagonist; trauma; gene therapy; cardiovascular disorder; endothelial disorder; cancer; anglogenic disorder; cardiac hypertrophy; atherosclerosis; hypertension; age-related macular degeneration; arterial restenosis; angina; rheumatoid arthritis; myocardial infarction; thrombophlebitis;
                                                                                                                                                                               Novel isolated interleukin 22 polypeptide useful for identifying IL-22 agonists and antagonists that are used for treating acute pancreatitis, chronic pancreatitis, pancreatic carcinoma
                                               Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
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                                             Maruoka EM,
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100.0%; Pred. No. 2.7e-86;
ive 0; Mismatches 0;
                                                 Gurney AL,
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                                          Goddard A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABB84993 standard; Protein; 179 AA
                                                                                                                                                                                                                                                                      Claim 11; Fig 2; 94pp; English.
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                                             Aggarwal S, Foster JS,
(GETH ) GENENTECH INC.
                                                                                                               WPI; 2002-280940/32.
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ABL88072 to ABL88258 encode the PRO proteins given in ABB84817 to ABB85032. The PRO proteins and polynucleotides have cardiant, cytostatic, antiangiogenic, hypotensive, vulnerary and antiarteriosclerotic activities, and can be used in gene therapy. The PRO polynucleotides, proteins, agonists and antagonists are useful for treating or diagnosing a cardiovascular, endothelial or engiogenic disorder in a mammal, e.g. cardiac hypertrophy, trauma, cancer, age-related macular degeneration, atherosclerosis, hypertension, arterial restenosis, rheumatoid arthritis, angina, myccardial infarctions, thrombophlebitis, lymphangitis, tumour angiogenesis (such as breast carcinoma and liver carcinoma) and wound healing. The PRO polynucleotides have applications
lymphangitis; tumour angiogenesis; breast carcinoma; liver carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gerber H, Gerritsen ME, Goddard A;
L, Hillan KJ, Marsters SA, Pan J, Paoni NF;
CK, Williams PM, Wood WI, Ye W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 One hundred and eighty seven nucleic acids encoding PRO polypeptides, useful in diagnosis and treatment of cardiovascular (e.g. myocardial infarction), endothelial or angiogenic disorders in a mammal
              wound healing; chromosome mapping; gene mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 11; Fig 354; 565pp; English.
                                                                                                                                                                                                                                                                                                                      2000US-242922P.
2000US-0709238.
2000WO-US30952.
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2001US-0828366.
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Stephan JF, Watanabe CK,
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                                                                     WO200200690-A2.
                                            Homo sapiens.
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05-APR-2001;
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in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping. ABL88259 to ABL88267 represent primers and probes used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           derived inducible factors (TIFs) also known as interleukin-21 (IL-21). TIF polymucleotides are upregulated by the cytokine, IL-9. IL-TIF or IL-21 molecules are implicated in activation of STAT transcription factors, acute phase proteins and inflammation. The present sequence is human TIF protein. The TIF gene is located on chromosome 12.
                                                                                                                                                                                                 1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRIFMLA
                                                                                                                                                                           1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stimulating expression of STAT transcription factor and inducing production of acute phase protein in a cell, involves contacting capable of expressing STAT with T cell derived inducible factors
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                                                                                                    100.0%; Score 903; DB 23;
100.0%; Pred. No. 2.7e-86;
iive 0; Mismatches 0;
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Best Local Similarity 100.0%; Pred. No. 2.7e-86;
Matches 179; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAE19237 standard; Protein; 179 AA.
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                                                                                                                                         Matches 179; Conservative
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                                                                      Sequence 179 AA;
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Search completed: December 5, 2003, 18:16:35 Job time : 61.9323 secs

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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/PCTUB_PUBCOMB.pep:*
6: /cgn2_6/ptodata/1/pubpaa/PCTUB_PUBCOMB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
8: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
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15: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
16: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
17: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
16: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
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17: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
18: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            684280 seqs, 185983659 residues
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Listing first 45 summaries
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903
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Perfect score:
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SUMMARIES	ΠD	US-09-728-911-15	US-09-87	96-60-SD	US-10-06	US-10-21	US-09-92	US-10-25	US-09-74	US-10-06	118-10-06	175-10-06	0.7.00	OS-10-06	US-10-06	113-10-06		90-01-90
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## ALIGNMENTS

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61 KEASLADINITDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
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100.0%; Pred. No. 3.1e-90;
Live 0; Mismatches 0;
                                                                                          JOHNSON THONORMATION:
APPLICANT: Xu, Wenfeng
TITLE OF INVENTION: Human Cytokine Receptor
FILE REPERBENCE: 99-93
CURRENT APPLICATION NUMBER: US/09/728,911
CURRENT FILING DATE: 2000-12-01
PRIOR PHILOR DATE: 1999-12-03
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 2000-09-13
PRIOR APPLICATION NUMBER: US 60/244,610
PRIOR FILING DATE: 2000-10-31
PRIOR FILING DATE: 2000-10-31
PRIOR FILING DATE: PRIOR WINDOWS Version 3.0
SEQ ID NO 15
ILENGTH: 179
TYPE: PRT
ORGANISM: Homo sapiens
US-09-728-911-15
                          Sequence 15, Application US/09728911
Patent No. US20022012669A1
GENERAL INFORMATION:
APPLICANT: Presnell, Scott R.
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Best Local Similarity 100.
Matches 179; Conservative
US-09-728-911-15
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PRIOR FILING DATE: 1999-05-19
PRIOR PAPLICATION NUMBER: 60/144,270
PRIOR PILING DATE: 1999-07-15
PRIOR APPLICATION NUMBER: 60/146,700
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-07-30
PRIOR PILING DATE: 1999-10-04
NUMBER OF SEQ ID NOS: 55
SOFTWARE: PERL PROGRAM
SEQ ID NO 18
LENGTH: 179
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Grimaldi, Christopher J.
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Goddard, Audrey
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Matches 179; Conservative
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APPLICANT: wood, William I.
TITLE OF INVENTION: INTERLEGION 2 POLYPEPTIDES, NUCLEIC ACIDS ENCODING
TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDERS
TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDERS
TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDERS
CURRENT APPLICATION NUMBER: US 60/169,495
PRIOR APPLICATION NUMBER: PCT/USO0/14042
PRIOR FILING DATE: 1999-12-07
PRIOR FILING DATE: 2000-08-22
PRIOR FILING DATE: 2000-08-24
PRIOR FILING DATE: 2000-08-24
PRIOR FILING DATE: 2000-08-24
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                               121 FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 903; DB 10; Length 179; 100.0%; Pred. No. 3.1e-90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: INCIDENT TOM APPLICANT: YUR, Henry APPLICANT: YUE, Henry APPLICANT: YUE, Henry APPLICANT: BANDWAN, Olga APPLICANT: BANDWAN, Olga APPLICANT: BANDWAN, Olga APPLICANT: AZINZAL, Yalda APPLICANT: AZINZAL, Yalda APPLICANT: LU, DAVID AIDA M. APPLICANT: LU, DAVID AIDA M. APPLICANT: PROPORTION: EXTRACELLULAR SIGNALING MOLECULES ITELE REPRENCE: PF-0701 USA CURRENT APPLICATION NUMBER: US/09/965,528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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Publication No. US20020187523A1
GENERAL INFORMATION:
                                                                                                                                                                                     Sequence 2, Application US/09870574
Patent No. US20020102723A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: INCYTE GENOMICS, INC.
                                                                                                                                                                                                                                                                              APPLICANT: Aggarwal, Sudeepta
APPLICANT: Xie, Ming-Hong
APPLICANT: Maruoka, Ellen M.
                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Foster, Jessica S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best_Local Similarity 100.0
Matches 179; Conservative
                                                                                                                                                                                                                                   GENERAL INFORMATION: APPLICANT: Gurney, Austin L.
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SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo Sapien
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                                                                                                                                                                                                                                                                          1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA 60
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APPLICANT: Gurney, Austin L.
APPLICANT: Weatababe, Colin K.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3230R1C1
CURRENT APPLICATION NUMBER: US/10/063,735
CURRENT APPLICATION NUMBER: US/10/063,735
ETIOL Application removed - See Palm or File Wrapper
NUMBER OF SEQ ID NOS: 170
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                                                                                                                                              Length 179;
NAME/KEY: misc feature; OTHER INFORMATION: Incyte ID No. US20020187523A1 5571181CD1
US-09-965-528-18
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                                                                                                                                          100.0%; Score 903; DB 10;
100.0%; Pred. No. 3.1e-90;
tive 0; Mismatches 0;
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APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Watanabe, Colin L.
TITLE OF INVENTION: SECRETED AND TRANSMENERANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE SPERENCE: P3530PLC3
CURRENT APPLICATION NUMBER: US/10/15,163
CURRENT PILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 60/05913
PRIOR PILING DATE: 1997-09-17
PRIOR PILING DATE: 1997-10-17
PRIOR PILING DATE: 1997-10-17
PRIOR PILING DATE: 1997-10-17
PRIOR PRILING DATE: 1997-10-17
PRIOR PILING DATE: 1997-10-24
PRIOR PILING DATE: 1997-10-24
PRIOR PILING DATE: 1997-12-17
PRIOR PILING DATE: 1998-03-22
PRIOR APPLICATION NUMBER: 60/05994
PRIOR PILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/07928
PRIOR PILING DATE: 1998-03-26
PRIOR PILING DATE: 1998-03-26
PRIOR PILING DATE: 1998-03-26
PRIOR PILING DATE: 1998-03-27
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Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 246
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                                                                                                                                                                                                        NGS-10-216-163-244

Sequence 244, Application US/10216163
FUBLICATION NO. US20030149239A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Pul J.
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin L.
APPLICANT: Smith, Victoria
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US-10-216-163-244
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LENGTH: 179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Gaps
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APPLICANT: Jacobs, Kenneth
APPLICANT: Pittman, Debra
APPLICANT: Fouser, Lynette
APPLICANT: Spaniding, Vikki
APPLICANT: Xuan, Defun
TITLE OF INVENTION: Composition and Method for Treating Inflammatory
FILE REPERENCE: GISSS CIP
CURRENT APPLICATION NUMBER: US/10/256,977
PRIOR APPLICATION NUMBER: US/10/084,298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 903; DB 12; Length 179; Best Local Similarity 100.0%; Pred. No. 3.1e-90; Matches 179; Conservative 0; Mismatches 0; Indels 0;
Sequence 8, Application US/09925055D
| Sequence 8, Application US/09925055D
| Publication No. US2030157096A1
| GENERAL INFORMATION:
| APPLICANT: Kindsvogel, Wayne R.
| APPLICANT: Topouzis, Stavros
| TITLE OF INVENTION: SOLUBLE ZCYTORII CYTOKINE RECEPTORS
| TITLE OF INVENTION: SOLUBLE ZCYTORII CYTOKINE RECEPTORS
| FILE REFREENCE: 00-56
| CURRENT APPLICATION NUMBER: US/09/925,055D
| CURRENT FILING DATE: 2000-08
| PRIOR APPLICATION NUMBER: US 60/223,827
| PRIOR FILING DATE: 2000-12-01
| NUMBER OF SEQ ID NOS: 35
| SOFTWARE: FastSEQ for Windows Version 3.0
| SEQ ID NO 8
| LEATH: 179
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PRIOR PLILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/270,823
PRIOR PLILING DATE: 2001-02-23
PRIOR PELING DATE: 2001-04-03
PRIOR PILING DATE: 2001-04-03
PRIOR FILING DATE: 1999-04-28
PRIOR FILING DATE: 1999-04-28
PRIOR PILING DATE: 1000-04-28
PRIOR SPELICATION NUMBER: 09/561,811
PRIOR PILING DATE: 2000-04-28
NUMBER OF SEQ ID NOS: 10
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US-10-256-977-2
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ORGANISM: homo sapiens
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LENGTH: 179
TYPE: PRT
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                                                                                                               1 MAALQKSVSSFIMGTLATSCILLIALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA 60
                                                                                                                                                                                                                                                                                                                                121 FLARLSNRLSTCHIBGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
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                                                                            1 MAALQKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA 60
                                                                                                                                                                                                                                                                                                   121 FLARLSNRLSTCHIEGDDLHIQRNVQKLKDTVKKLGESGEIKAIGELDLLFMSLRNACI 179
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APPLICANT: Goddard, Audrey
APPLICANT: Goddwski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
APPLICANT: WOOD, WILL SERRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3230R1C1
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                             0; Gaps
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                             Indels
                             0
Pred. No. 3.1e-90;
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TITLE OF INVENTION: NOVEL CYTOKINE ZCYTO18
FILE REFRENCE: 99-106
CURRENT PELLING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: US 60/172,105
PRIOR APPLICATION NUMBER: US 60/172,105
PRIOR APPLICATION NUMBER: US 60/172,105
PRIOR APPLICATION NUMBER: US 60/74*,***
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 2
LENGTH. 179
                             0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               %2-09-746-375-2
; Sequence 2, Application US/09746375
; Publication No. US20030170823A1
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
  100.08;
                             Matches 179; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 179; Conservative
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US-09-746-375-2
  Best Local Similarity
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61 KEASLADINITDVRLIGEKLFHGVSMSERCYLMKQVLNFTLEEVLFPQSDRFQPYMQEVVP 120
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                             Length 179;
                                                                                                                                                                                                                                        Indels
              CURRENT FILING DATE: 2002-05-02
Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 170
SEQ ID NO 154
LENGTH: 179
                                                                                                                                                                                               100.0%; Score 903; DB 12;
100.0%; Pred. No. 3.1e-90;
ative 0; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/10/066,198
CURRENT FILING DATE: 2002-02-01
CURRENT APPLICATION NUMBER: US/10/063,526
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PRIOR FILING DATE: 2001-11-15
PRIOR PELING DATE: 2001-11-15
PRIOR PELING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/05914
PRIOR APPLICATION NUMBER: 60/05915
PRIOR PILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-17
PRIOR PLICATION NUMBER: 60/05958
PRIOR PLICATION NUMBER: 60/05958
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Sherman Pong
Wei-Qiang Gao
Hanspeter Gerber
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Daniel Tumas
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P.Mickey Williams
William I. Wood
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Austin L. Gurney
Ivar J. Kljavin
Jennie P. Mather
Mary A. Napier
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APPLICANT: Kevin P. Baker
APPLICANT: David A. Botstein
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Margaret Ann Roy
                                                                                                                                                                                               Query Match
Best Local Similarity 100.0
Matches 179; Conservative
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Audrey Goddard
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Dan L. Eaton
                                                                                                                     TYPE: PRT
ORGANISM: Homo Sapien
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PRIOR PELING DATE: 1998-10.09

PRIOR APPLICATION NUMBER: 09/24311

PRIOR APPLICATION NUMBER: 09/254465

PRIOR APPLICATION NUMBER: 09/254465

PRIOR APPLICATION NUMBER: 09/254465

PRIOR PLING DATE: 1999-06-14

PRIOR PLING DATE: 1999-06-15

PRIOR PLING DATE: 1999-06-14

PRIOR PLING DATE: 1999-06-25

PRIOR PLING DATE: 1999-10-18

PRIOR PLING DATE: 2000-09-18

PRIOR PLING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: 09/8663

PRIOR PLING DATE: 2001-05-30

PRIOR PLING DATE: 1998-09-17

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE REFERENCE: P3230R1C1
CURRENT APPLICATION NUMBER: US/10/063,586
CURRENT FILING DATE: 2002-05-03
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 170
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                                                                                                                                                                                                                                                                                            100.0%; Score 903; DB 12; Length 179; 100.0%; Pred. No. 3.1e-90;
                                                                                                                                                                                                                                                                                                                                               0; Mismatches
PRIOR FILING DATE: 1999-03-06
PRIOR APPLICATION NUMBER: PCT/US99/12252
PRIOR FILING DATE: 1999-06-02
PRIOR FILING DATE: 1999-09-01
PRIOR FILING DATE: 1999-09-01
PRIOR PELING DATE: 1999-09-01
PRIOR PELING DATE: 1999-09-08
PRIOR FILING DATE: 1999-09-08
PRIOR PILING DATE: 1999-09-08
PRIOR PILING DATE: 1999-09-15
PRIOR PLING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/21090
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APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney Austin L.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
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Gerritsen, Mary E.
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Matches 179; Conservative
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US-10-063-586-154
                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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US-10-063-586-154
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APPLICANT:
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                                                                                                                                                                         APPLICANT: Elivaroff, Ellen
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddward, Audrey
APPLICANT: Goddward, Audrey
APPLICANT: Goddward, Lean J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Granele, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICAN
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Best Local Similarity 100.0%; Pred. No. 3.1e-90;
Matches 179; Conservative 0; Mismatches 0; Indels 0
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Sequence 154, Application US/10063510 Publication No. US20030180837A1 GENERAL INFORMATION:
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Grimaldi, Christopher J.
Gurney, Austin L.
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APPLICANT: Gerriteen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher
APPLICANT: Grimsey, Marstin L.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
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APPLICANT: Filvaroff, Blen
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
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ORGANISM: Homo Sapien
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US-10-063-514-154
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                   1 MAALOKSVSSFLMGTLATSCLLLLALLVQGGAAAPISSHCRLDKSNFQQPYITNRTFMLA 60
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APPLICANT: Goddard, Audray
APPLICANT: Goddard, Audray
APPLICANT: Goddowski, Paul J.
APPLICANT: Gurmaldi, Christopher J.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANB POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REPERBURCE: P23.30FRC1
CURRENT APPLICATION NUMBER: 105.10/063,516
CURRENT FILING DATE: 2002-05-01
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Publication No. US20030181636A1
GENERAL INFORMATION:
APPLICANT: Baton, Dan L.
APPLICANT: Filvaroff, Ellen
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APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
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APPLICANT: Filvaroff, Ellen
APPLICANT: Gerritsen, Mary E.
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Goddard, Audrey
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Best Local Similarity 100.03
Matches 179; Conservative
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ORGANISM: Homo Sapien
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APPLICANT:
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                     FILE REFERENCE: P3230R1C1
CURRENT APPLICATION NUMBER: US/10/063,523
CURRENT FILING DATE: 2002-05-02
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 170
SEQ ID NO 154
ACIDS ENCODING THE SAME
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Job time : 346.883 secs
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Matches 179; Conservative
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ORGANISM: Homo Sapien
  TITLE OF INVENTION:
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